

Replaced

Recd PTO

20 APR 2001

SEQUENCE LISTING

<110> KIMCHI, Adi

<120> DAP-KINASE RELATED PROTEIN

<130> KIMCHI2A

<140> 09/719,748

<141> 2000-12-15

<150> 60/089,294

<151> 1998-06-15

<160> 14

<170> PatentIn Ver. 2.0

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Glu Glu Leu Gly Ser Gly Gln Phe Ala Ile Val Lys Lys Cys Arg Glu

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aag agc acg ggg ctt gag tat gca gcc aag ttc atc aag aag cgg cag 205

Lys Ser Thr Gly Leu Glu Tyr Ala Ala Lys Phe Ile Lys Lys Arg Gln

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agc cgg gcg agc cgg cgc ggt gtg agc cgg gag gag atc gag cgg gag 253

Ser Arg Ala Ser Arg Arg Gly Val Ser Arg Glu Glu Ile Glu Arg Glu

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gtg agc atc ctg cgg cag gtg ctg cac cac aat gtc atc acg ctg cac 301

Val Ser Ile Leu Arg Gln Val Leu His His Asn Val Ile Thr Leu His

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gac gtc tat gag aac cgc acc gac gtg gtg cac atc ctt gag cta gtg 349

Asp Val Tyr Glu Asn Arg Thr Asp Val Val His Ile Leu Glu Leu Val

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tct gga gga gag ctc ttc gat ttc ctg gcc cag aag gag tca ctg agt 397

Ser Gly Gly Glu Leu Phe Asp Phe Leu Ala Gln Lys Glu Ser Leu Ser

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105

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tac ctt cac aca aag aaa att gct cac ttt gat ctc aag cca gaa aac	493
Tyr Leu His Thr Lys Lys Ile Ala His Phe Asp Leu Lys Pro Glu Asn	
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att atg ttg tta gac aag aat att ccc att cca cac atc aag ctg att	541
Ile Met Leu Leu Asp Lys Asn Ile Pro Ile Pro His Ile Lys Leu Ile	
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gac ttt ggt ctg gct cac gaa ata gaa gat gga gtt gaa ttt aag aat	589
Asp Phe Gly Leu Ala His Glu Ile Glu Asp Gly Val Glu Phe Lys Asn	
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Ile Leu Leu Ser Gly Ala Ser Pro Phe Leu Gly Asp Thr Lys Gln Glu	
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Thr Leu Ala Asn Ile Thr Ser Val Ser Tyr Asp Phe Asp Glu Glu Phe	
225 230 235 240	
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Phe Ser His Thr Ser Glu Leu Ala Lys Asp Phe Ile Arg Lys Leu Leu	
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Val Lys Glu Thr Arg Lys Arg Leu Thr Ile Gln Glu Ala Leu Arg His	
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Pro Trp Ile Thr Pro Val Asp Asn Gln Gln Ala Met Val Arg Arg Glu	
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tct gtg gtc aat ctg gag aac ttc agg aag cag tat gtc cgc agg cgg	973
Ser Val Val Asn Leu Glu Asn Phe Arg Lys Gln Tyr Val Arg Arg Arg	
290 295 300	
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Trp Lys Leu Ser Phe Ser Ile Val Ser Leu Cys Asn His Leu Thr Arg	
305 310 315 320	
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Ser Leu Met Lys Lys Val His Leu Arg Pro Asp Glu Asp Leu Arg Asn	
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 35 40 45
 Ser Arg Ala Ser Arg Arg Gly Val Ser Arg Glu Glu Ile Glu Arg Glu
 50 55 60
 Val Ser Ile Leu Arg Gln Val Leu His His Asn Val Ile Thr Leu His
 65 70 75 80
 Asp Val Tyr Glu Asn Arg Thr Asp Val Val His Ile Leu Glu Leu Val
 85 90 95
 Ser Gly Gly Glu Leu Phe Asp Phe Leu Ala Gln Lys Glu Ser Leu Ser
 100 105 110
 Glu Glu Glu Ala Thr Ser Phe Ile Lys Gln Ile Leu Asp Gly Val Asn
 115 120 125

Tyr Leu His Thr Lys Lys Ile Ala His Phe Asp Leu Lys Pro Glu Asn
 130 135 140
 Ile Met Leu Leu Asp Lys Asn Ile Pro Ile Pro His Ile Lys Leu Ile
 145 150 155 160
 Asp Phe Gly Leu Ala His Glu Ile Glu Asp Gly Val Glu Phe Lys Asn
 165 170 175
 Ile Phe Gly Thr Pro Glu Phe Val Ala Pro Glu Ile Val Asn Tyr Glu
 180 185 190
 Pro Leu Gly Leu Glu Ala Asp Met Trp Ser Ile Gly Val Ile Thr Tyr
 195 200 205
 Ile Leu Leu Ser Gly Ala Ser Pro Phe Leu Gly Asp Thr Lys Gln Glu
 210 215 220
 Thr Leu Ala Asn Ile Thr Ser Val Ser Tyr Asp Phe Asp Glu Glu Phe
 225 230 235 240
 Phe Ser His Thr Ser Glu Leu Ala Lys Asp Phe Ile Arg Lys Leu Leu
 245 250 255
 Val Lys Glu Thr Arg Lys Arg Leu Thr Ile Gln Glu Ala Leu Arg His
 260 265 270
 Pro Trp Ile Thr Pro Val Asp Asn Gln Gln Ala Met Val Arg Arg Glu
 275 280 285
 Ser Val Val Asn Leu Glu Asn Phe Arg Lys Gln Tyr Val Arg Arg Arg
 290 295 300
 Trp Lys Leu Ser Phe Ser Ile Val Ser Leu Cys Asn His Leu Thr Arg
 305 310 315 320
 Ser Leu Met Lys Lys Val His Leu Arg Pro Asp Glu Asp Leu Arg Asn
 325 330 335
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 35 40 45
 Ile Glu Arg Glu Val Ser Ile Leu Lys Glu Ile Gln His Pro Asn Val
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 Ile Thr Leu His Glu Val Tyr Glu Asn Lys Thr Asp Val Ile Leu Ile
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 Leu Glu Leu Val Ala Gly Gly Glu Leu Phe Asp Phe Leu Ala Glu Lys
 85 90 95
 Glu Ser Leu Thr Glu Glu Glu Ala Thr Glu Phe Leu Lys Gln Ile Leu
 100 105 110
 Asn Gly Val Tyr Tyr Leu His Ser Leu Gln Ile Ala His Phe Asp Leu
 115 120 125
 Lys Pro Glu Asn Ile Met Leu Leu Asp Arg Asn Val Pro Lys Pro Arg
 130 135 140
 Ile Lys Ile Ile Asp Phe Gly Leu Ala His Lys Ile Asp Phe Gly Asn
 145 150 155 160
 Glu Phe Lys Asn Ile Phe Gly Thr Pro Glu Phe Val Ala Pro Glu Ile
 165 170 175
 Val Asn Tyr Glu Pro Leu Gly Leu Glu Ala Asp Met Trp Ser Ile Gly
 180 185 190
 Val Ile Thr Tyr Ile Leu Leu Ser Gly Ala Ser Pro Phe Leu Gly Asp
 195 200 205
 Thr Lys Gln Glu Thr Leu Ala Asn Val Ser Ala Val Asn Tyr Glu Phe
 210 215 220
 Glu Asp Glu Tyr Phe Ser Asn Thr Ser Ala Leu Ala Lys Asp Phe Ile
 225 230 235 240
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 Ser Leu Gln His Pro Trp Ile
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 35 40 45
 Ile Glu Arg Glu Val Asn Ile Leu Arg Glu Ile Arg His Pro Asn Ile
 50 55 60
 Ile Thr Leu His Asp Ile Phe Glu Asn Lys Thr Asp Val Val Leu Ile
 65 70 75 80
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 85 90 95
 Glu Ser Leu Thr Glu Asp Glu Ala Thr Gln Phe Leu Lys Gln Ile Leu
 100 105 110
 Asp Gly Val His Tyr Leu His Ser Lys Arg Ile Ala His Phe Asp Leu
 115 120 125
 Lys Pro Glu Asn Ile Met Leu Leu Asp Lys Asn Val Pro Asn Pro Arg
 130 135 140
 Ile Lys Leu Ile Asp Phe Gly Ile Ala His Lys Ile Glu Ala Gly Asn
 145 150 155 160
 Glu Phe Lys Asn Ile Phe Gly Thr Pro Glu Phe Val Ala Pro Glu Ile
 165 170 175
 Val Asn Tyr Glu Pro Leu Gly Leu Glu Ala Asp Met Trp Ser Ile Gly
 180 185 190
 Val Ile Thr Tyr Ile Leu Leu Ser Gly Ala Ser Pro Phe Leu Gly Glu
 195 200 205
 Thr Lys Gln Glu Thr Leu Thr Asn Ile Ser Ala Val Asn Tyr Asp Phe
 210 215 220
 Asp Glu Glu Tyr Phe Ser Asn Thr Ser Glu Leu Ala Lys Asp Phe Ile
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 Arg Arg Leu Leu Val Lys Asp Pro Lys Arg Arg Met Thr Ile Ala Gln
 245 250 255
 Ser Leu Glu His Ser Trp Ile
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His	Glu	Val	Tyr	Glu	Thr	Ala	Ser	Glu	Met	Ile	Leu	Val	Leu	Glu	Tyr
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Ala	Ala	Gly	Gly	Glu	Ile	Phe	Asp	Gln	Cys	Val	Ala	Asp	Arg	Glu	Glu
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130						135				140					
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145				150						155				160	
Arg	Glu	Ile	Met	Gly	Thr	Pro	Glu	Tyr	Val	Ala	Pro	Glu	Ile	Leu	Ser
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Thr	Tyr	Val	Met	Leu	Thr	Gly	Ile	Ser	Pro	Phe	Leu	Gly	Asn	Asp	Lys
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Leu	Leu	Val	Lys	Lys	Pro	Glu	Asp	Arg	Ala	Thr	Ala	Glu	Glu	Cys	Leu
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Lys Lys Arg Arg Arg Gly Gln Asp Cys Arg Ala Glu Ile Leu His Glu
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Ile Ala Val Leu Glu Leu Ala Lys Ser Cys Pro Arg Val Ile Asn Leu
50 55 60

His Glu Val Tyr Glu Asn Thr Ser Glu Ile Ile Leu Ile Leu Glu Tyr
65 70 75 80

Ala Ala Gly Gly Glu Ile Phe Ser Leu Cys Leu Pro Glu Leu Ala Glu
85 90 95

Met Val Ser Glu Asn Asp Val Ile Arg Leu Ile Lys Gln Ile Leu Glu
100 105 110

Gly Val Tyr Tyr Leu His Gln Asn Asn Ile Val His Leu Asp Leu Lys
115 120 125

Pro Gln Asn Ile Leu Leu Ser Ser Ile Tyr Pro Leu Gly Asp Ile Lys
130 135 140

Ile Val Asp Phe Gly Met Ser Arg Lys Ile Gly His Ala Cys Glu Leu
145 150 155 160

Arg Glu Ile Met Gly Thr Pro Glu Tyr Leu Ala Pro Glu Ile Leu Asn
165 170 175

Tyr Asp Pro Ile Thr Thr Ala Thr Asp Met Trp Asn Ile Gly Ile Ile
180 185 190

Ala Tyr Met Leu Leu Thr His Thr Ser Pro Phe Val Gly Glu Asp Asn
195 200 205

Gln Glu Thr Tyr Leu Asn Ile Ser Gln Val Asn Val Asp Tyr Ser Glu
210 215 220

Glu Thr Phe Ser Ser Val Ser Gln Leu Ala Thr Asp Phe Ile Gln Ser
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Ser His Ser Trp Leu
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